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Discovery of selective 2,4-diaminopyrimidinebased photoaffinity probes for glyoxalase I⁺

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Glyoxalase I (GLO-1) plays a critical role in the detoxification of 2-oxoaldehydes and is highly expressed in cancer cells. Through photo-affinity labelling and affinity pull-down approaches, a series of 2,4diaminopyrimidine compounds were discovered to selectively bind to GLO-1 in cells. These compounds show potent inhibition of GLO-1 enzyme activity and prevent proliferation of cancer cells. The cell permeable and "clickable" photoaffinity probe L1-Bpyne presented here could be a valuable tool for profiling GLO-1 in live cells.

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Introduction

The glyoxalase system, composed of glyoxalase I (GLO-1, EC 4.1.1.5) and glyoxalase II (GLO-2, EC 3.1.2.6), is an important

20 and ubiquitous detoxification component of cellular metabolism in most cells.¹ Methylglyoxal (MG), a by-product in glycolysis, is cytotoxic due to its covalent binding to DNA and proteins in cells.^{1,2} Newly produced MG reacts with glutathione (GSH), an abundant nucleophile in cells, to form hemithioacetal. GLO-1 25 catalyses the isomerisation of hemithioacetal into S-D-lactoylglutathione. Subsequently GLO-2 catalyses the conversion of lactoylglutathione into lactic acid and GSH via hydrolysis.3 The glyoxalase system is highly expressed in cancer and inflammatory cells to rapidly remove the cytotoxic MG.⁴

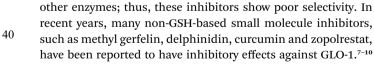
Overexpression of GLO-1 has been associated with multi-

drug resistance in cancer chemotherapy.5 Much effort has been made to develop inhibitors of GLO-1 as an anti-cancer therapy. Most of the early GLO-1 inhibitors are GSH deriva-

tives, but this class of compounds has poor pharmacokinetic properties, as GSH remains charged under physiological conditions.⁶ Additionally, GSH is a substrate or cofactor of many

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† Electronic supplementary information (ESI) available: Detailed synthetic 50 procedures, ¹H and ¹³C NMR spectra, HPLC chromatograms for compounds L1, L2, L3, L1-Biotin and L1-Bpyne; detailed procedures for biological assays; mass spectroscopy data for GLO-1. See DOI: 10.1039/c3md00286a

Despite advancements in the development of GLO-1 inhibitors, a selective GLO-1 labelling probe has not been reported.

The 2,4-diaminopyrimidine scaffold has been widely used in pharmaceutical agents, including the dihydrofolate reductase 20 inhibitor pyrimethamine, and the antibiotics iclaprim and trimethoprim.¹¹ Recently, 2,4-diaminopyrimidine also served as a common template to generate inhibitors for a number of kinases, including c-Met,12 MK2,13 ALK,14 and SYK,15 Since 2011, several research groups have developed potent LRRK2 inhibi-25 tors based on the 2,4-diaminopyrimidine scaffold.¹⁶⁻¹⁸ Of these compounds, CZC-25146 was found to attenuate Parkinson's disease-related toxicity in human neurons (Fig. 1A). While the in vitro selectivity of this type of kinase inhibitor can be assessed using various techniques,¹⁹⁻²¹ the selectivity of these 30 compounds in more complex biological systems is still unknown.²² To examine the selectivity of this type of inhibitors

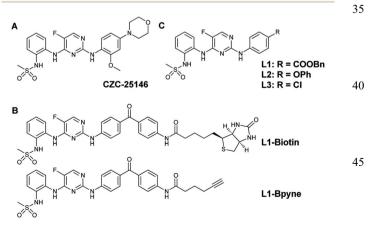


Fig. 1 Chemical structures of (A) the 2,4-diaminopyrimidine-based 50 kinase inhibitor CZC-25146; (B) the photoaffinity probes L1-Biotin and L1-Bpyne; and (C) the analogues of the 2,4-diaminopyrimidine series compounds used in this study.

in mammalian cells, we synthesised two photoaffinity probes, **L1-Biotin** and **L1-Bpyne**, to detect these compounds cellular targets (Fig. 1B). As a result, GLO-1 was identified as a major target of these 2,4-diaminopyrimidine compounds in live cells.

5 Several synthetic analogues of CZC-25146 were further tested in *in vitro* enzyme activity and cell proliferation assays (Fig. 1C). These compounds inhibited the enzyme activity of GLO-1 and cancer cell proliferation, and the potency of these molecules depended on the glucose concentrations in culture media.
10 Finally, these compounds were shown to cause MG accumulation in live cells and induce cell apoptosis.

Results and discussion

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First, we examined the potential targets of 2,4-diaminopyrimidine compounds in cells and designed a biotinylated photoaffinity probe L1-Biotin to label and pull-down its binding proteins in the cellular proteome (Fig. 1B). A benzophenone group was introduced to covalently link the probe and its 20 potential targets upon photo irradiation. Human embryo kidney 293T (HEK293T) cell lysate was pre-treated with streptavidin-sepharose beads to reduce endogenously biotinylated proteins. The cells were then incubated with 0.1 or 1 µM L1-25 Biotin with and without 50 µM compound L1 overnight at 4 °C. After photo-crosslinking by irradiation at 365 nm for 1 hour on ice, the L1-Biotin binding proteins were separated by SDS-PAGE and detected by streptavidin-HRP (Fig. 2A). Alternatively, biotinylated proteins in lysates, both endogenous and those bound 30 to the L1-Biotin probe, were first enriched with streptavidin-

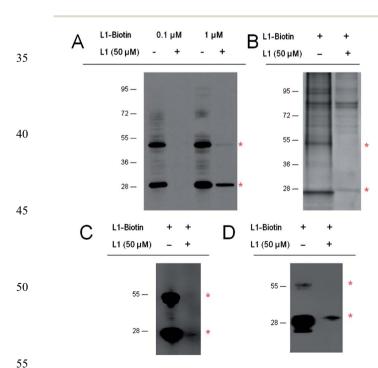


Fig. 2 (A) HEK293T cell lysate labelling by L1-Biotin with and without competitor L1; (B) silver staining of L1-Biotin binding proteins enriched by streptavidin–sepharose beads; (C) western blot validation with GLO-1 antibody; (D) recombinant glyoxalse-1 protein labelling by L1-Biotin.

sepharose beads, then separated by SDS-PAGE and visualised by 1 silver staining (Fig. 2B). In both experiments, the L1-Biotin probe bound specifically to two protein bands at approximately 28 kDa and 55 kDa. The two corresponding bands in silver staining gels were excised, destained, digested by trypsin and 5 analysed by MALDI-TOF mass spectrometry. Both bands were identified as human glyoxalase I (see ESI⁺). As the literature indicates, GLO-1 exists in both monomeric and dimeric forms in cells;^{2,23} the higher molecular weight band could represent the GLO-1 dimer, and the lower molecular weight band could 10 represent the monomer. The appearance of the dimeric form band at 55 kDa in the SDS-PAGE experiments might be caused by our probe molecules and cellular modifications in HEK293T cells. These results were further confirmed by western blotting 15 using glyoxalase I antibody (Fig. 2C). Finally, recombinant human GLO-1 protein was also successfully labelled by L1-Biotin (Fig. 2D) in a separate experiment. Competitive binding is clearly observed using the structurally similar compound L1, suggesting that GLO-1 was specifically labelled by the probe L1-20 Biotin.

Further labelling experiments were performed to evaluate the detection range of the probe L1-Biotin and the effect of UV irradiation time. HEK293T lysates were incubated with different concentrations of L1-Biotin; 0.5 to 1 μ M of probe was sufficient to label GLO-1, whereas higher concentrations of the probe increased non-specific labelling (Fig. 3A). At a steady probe concentration at 0.5 μ M, cell lysates were incubated with L1-Biotin and irradiated with UV light for various periods of time. The band intensity increased with the UV irradiation time and became saturated after approximately 1 hour (Fig. 3B).

After GLO-1 was identified as a potential target in cell lysates, we attempted to confirm it in live cells. However, the biotinylated photoaffinity probe **L1-Biotin** was unable to bind to proteins in live cells (data not shown), likely due to the low 35 trans-membrane permeability of the biotin tag. We then designed the probe **L1-Bpyne** for labelling studies of GLO-1 in live cells (Fig. 1B). Bio-orthogonal conjugation reactions have been successfully employed to identify covalent targets of small molecules in a variety of biological systems. The copper(I) catalysed [3 + 2] azide–alkyne cycloaddition (CuAAC) reaction is

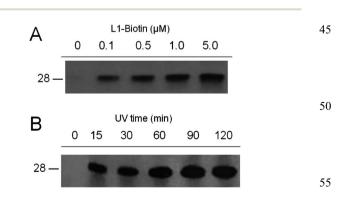


Fig. 3 (A) HEK293T cell lysates treated with different concentrations of probe L1-Biotin; (B) HEK293T cell lysates treated with 0.5 μM of L1-Biotin and irradiated by long-wave UV light for different lengths of time.

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one of the most commonly used methods.²⁴ Thus, in the new **L1-Bpyne** probe, an alkyne tag replaced the biotin tag in **L1-Biotin** to facilitate cellular studies.

We then attempted to label GLO-1 using this new probe. For
 in vitro labelling, HEK293T lysates were incubated with 1 μM L1 Bpyne with and without excess compound L1. After UV irradiation and CuAAC conjugation with tetramethylrhodamine azide (TAMRA-N₃), the samples were analysed by SDS-PAGE and detected by in-gel fluorescence scan. For *in situ* labelling, live

- 10 HEK293T cells were incubated with 1 μ M of L1-Bpyne with and without excess compound L1. The cells were irradiated by UV for 1 hour at 37 °C and washed with ice-cold PBS buffer solution to remove unbound probes. Following centrifugation and homogenisation by lysis buffer, the samples were subjected to
- 15 the CuAAC reaction with the same conditions as those used for the *in vitro* labelling. Both monomeric and dimeric forms of GLO-1 were selectively labelled by the probe L1-Bpyne *in vitro* and *in situ*. The labelling showed specific binding competition by compound L1 (Fig. 4A). Encouraged by the results of *in situ*
- ²⁰ by compound L1 (Fg. H). Introduciged by the results of the state labelling, the intracellular CuAAC reaction with TAMRA-N₃ was performed to show the cellular localisation of L1-Bpyne targets. HeLa cells were cultured in normal culture medium and treated with 20 μM L1-Bpyne for 2 hours with or without 100 μM
 ²⁵ compound L1 as the competitor. Following UV irradiation, the cells were fixed, permeabilised, and subjected to CuAAC reaction with TAMRA-N₃ (Fig. 4B). Staining of the cells gave a strong signal as GLO-1 is a relatively abundant enzyme in cytosol. The labelling was significantly reduced following competitive binding by excess compound L1.

Because GLO-1 appeared as a major cellular target of 2,4diaminopyrimidine probes, we next investigated whether these types of compounds could inhibit the catalytic activity of GLO-1 *in vitro*. A spectrophotometric method was employed by monitoring the increase in UV absorbance at 240 nm from the formation of *S*-D-lactoylglutathione at 25 °C.²⁵ The **L1-Bpyne** probe was the most potent inhibitor with an IC₅₀ value of 26 nM, while other analogues **L1–L3** showed IC₅₀ values of

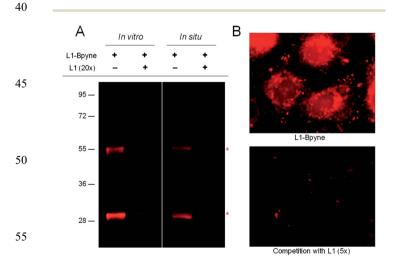


Fig. 4 (A) In vitro labelling of HEK293T cell lysates and in situ labelling

of live HEK293T cells by L1-Bpyne (1 $\mu M);$ (B) in-cell CuAAC reaction revealed the cellular distribution of L1-Bpyne.

approximately 50 nM (Table 1), which suggests that the benzophenone moiety and the alkyne tag did not significantly alter the binding preference of these 2,4-diaminopyrimide compounds towards GLO-1.

Proliferation assays were performed to examine the effects of 5 these compounds in cells. The primary cellular function of GLO-1 is MG detoxification, and the MG concentration has been reported to be elevated in high glucose conditions.^{25,26} Therefore, three types of cancer cells were cultured in Dulbecco's Modified Eagle Medium (DMEM) with two concentrations of 10 glucose at 5 mM (approximately the normal *in vivo* glucose level) and 25 mM (high glucose level). Cell viability was assessed using a CellTiter-Glo® luminescent kit after incubating for 48 hours with various concentrations of the tested compounds. These 15 compounds had GI₅₀ values at low to mid micromolar concentrations (Table 1). In accordance with the data of the enzymatic assay, the L1-Bpyne probe was also the most potent inhibitor in the cell proliferation assays. Between two human lung cancer cell lines, NCI-H522 cell, which was reported to express high 20 intrinsic GLO-1 activity, is more susceptible to the GLO-1 inhibition; A549 cell, which was reported to express relatively low intrinsic GLO-1 activity, showed about 5-fold less sensitivity towards L1-Bpyne.²⁷ HeLa cell, a human cervical cancer cell line, could also be potently inhibited by this series of inhibitors. The 25 glucose contents in the culture media had low to moderate effects on the inhibition of tumor cell proliferation by this series of compounds. As expected, high concentrations of glucose increased cellular sensitivity to GLO-1 inhibition. The GI₅₀ values of compounds L1-L3 increased up to 2.6-fold in normal 30 glucose medium compared to high glucose medium.

Because GLO-1 inhibition causes methylglyoxal accumulation followed by random DNA and protein labelling, we further investigated whether our compounds could induce an increase of the cellular MG concentration. HeLa cells were grown in 35 normal or high glucose media for 3 days followed by treatment with DMSO or 5 µM of L1-Bpyne for 24 hours. The cells were harvested, washed with PBS buffer, resuspended in distilled water and boiled. The MG content in the supernatant was 40 detected using the same spectrophotometric method as that in the aforementioned enzyme activity assay. The MG concentrations were slightly elevated in the high glucose medium with untreated cells but were almost doubled in the L1-Bpyne treated cells (Fig. 5A). This suggests that L1-Bpyne inhibited GLO-1 45 activity in HeLa cells. Cellular MG accumulation commenced upon GLO-1 inhibition.

Finally, we tested whether GLO-1 inhibition would cause cell apoptosis. HeLa cells were treated with various concentrations of **L1-Bpyne** for 24 hours, the degradation of the apoptosis marker protein caspase-3 clearly indicated cell apoptosis (Fig. 5B). Moreover, after HeLa cells were treated with 5μ M of **L1-Bpyne** for 24 hours, the cells were fixed with paraformaldehyde and stained with 4',6-diamidino-2-phenylindole (DAPI), the cell nuclei also appeared fragmented (Fig. 5C).

MG is a highly reactive α -oxoaldehyde that reacts predominantly with the lysine or arginine residues of proteins and the guanyl residues in DNA and RNA. It may cause abnormal enzyme activity and defects during transcription.²⁸ Because 50

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Table 1	Inhibition of GLO-1 enz	vme activity and cance	er cells proliferation b	y 2,4-diaminopyrimidine compounds ^a

Compound	GLO-1 (nM)	NCI-H522-HG (μM)	NCI-H522-NG (μM)	HeLa-HG (µM)	HeLa-NG (µM)	A549-HG (µM)	A549-NG (μM)
L1-Bpyne	26.4	2.9	2.7	4.6	5.7	13.3	15.5
L1	51.3	16.5	40.2	37.0	61.9	>50	>50
L2	45.8	7.3	8.3	7.3	14.0	26.9	37.4
L3	53.9	12.2	25.0	23.2	61.0	>50	>50

^{*a*} Values are averages of at least two independent runs. HG: high glucose level; NG: normal glucose level.

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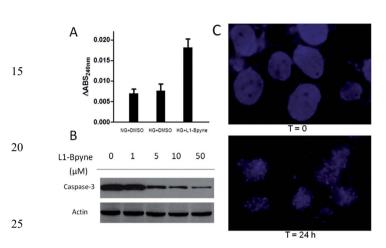


Fig. 5 (A) Intracellular MG concentration upon L1-Bpyne treatment; (B) HeLa cells were treated with different concentrations of L1-Bpyne and the intact caspase-3 levels were evaluated by western-blot; (C) HeLa cells were treated with 5 μ M of L1-Bpyne for 24 hours and stained by DAPI.

tumour tissue has higher glycolysis activity than normal
 tissue^{29,30} and the glyoxalase system is highly activated in cancer cells,^{31,32} GLO-1 inhibitors could be selectively cytotoxic towards cancer cells. Our compounds did show inhibition of proliferation of several human cancer cells, thus indicating their potential application to cancer therapy. The relative large

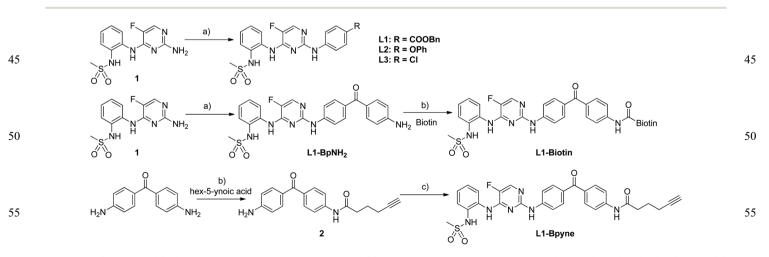
potency shift of these compounds from the enzymatic assay to the cell proliferation assay could result from multiple factors including physiochemical properties of compounds and the network linking GLO-1 activity to cell death. A combination of GLO-1 inhibitors with compounds targeting multiple pathways may be an attractive approach to achieve significant anti-cancer effects in the clinic.

Chemistry

The syntheses of compounds L1–L3, L1-Biotin and L1-Bpyne are depicted in Scheme 1. Each was readily prepared using literature procedures with slight modifications. The common precursor 1 was coupled with substituted anilines by an S_NAr 25 reaction to provide L1, L2, and L3 in good yields. L1-Biotin was prepared from the intermediate L1-BpNH₂ and biotin using standard coupling conditions for amide bond formation. Compounds 1 and 2 underwent a Hartwig–Buchwald amination reaction to furnish L1-Bpyne in good yield. Detailed synthetic procedures and compound characterisation data are provided in the ESI.[†]

Conclusions

In summary, we have discovered two photoaffinity probes, L1-**Biotin** and L1-Bpyne, which specifically label GLO-1 in cells. The biotinylated probe can specifically label and pull-down GLO-1 *in vitro*, while the alkyne probe is able to passively penetrate living



Scheme 1 Synthesis of GLO-1 inhibitors. *Reagents and conditions*: (a) substituted anilines, concentrated HCl, *n*-BuOH, 120 °C, 16–18 hours; (b) HATU, Hunig's base, DMF, 0 °C to rt, overnight; (c) 1, Pd₂(dba)₃, DPBP, K₂CO₃, nBuOH, 100 °C, 4 hours.

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- cells and label GLO-1 *in situ*. Furthermore, all compounds potently inhibited the enzyme activity of GLO-1 with IC_{50} values in the range of 26 to 54 nM. These compounds prevented cancer cell proliferation, increased intracellular levels of methyl-
- ⁵ glyoxal, and induced cell apoptosis. Interestingly, higher glucose content in the cell medium increased the cytotoxicity of these compounds. Taken together, this series of 2,4-diaminopyrimidine compounds directly interacts with GLO-1 in cells. Moreover, **L1-Bpyne** is both a potent inhibitor and
- 10 a selective probe of GLO-1, and could be a powerful tool in studies of this enzyme. With this cell permeable probe, GLO-1 was successfully labelled both *in vitro* and *in situ*. This probe could also be utilised to label GLO-1 in tissues of animal models, though the long-wave UV light used in the photo-
- crosslinking step could be harmful to surrounding tissues and limits the application of this probe in whole animal studies. Further research is underway to understand the binding mode of this type of probe, including the exact binding site and specific binding interactions with GLO-1, and its potential use
- in combination with other anti-cancer agents.

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Notes and references

45

- 35 1 P. Thornalley, *Biochem. J.*, 1990, **269**, 1–11.
 - 2 P. Thornalley, Biochem. Soc. Trans., 2003, 31, 1372-1377.
 - 3 F. Himo and Per E. M. Siegbahn, *J. Am. Chem. Soc.*, 2001, **123**, 10280–10289.
 - 4 R. Vince and S. Daluge, J. Med. Chem., 1971, 14, 35-37.
- 5 D. Creighton, Z. Zheng, R. Holewinski, D. Hamilton and J. Eiseman, *Biochem. Soc. Trans.*, 2003, 31, 1378–1382.
 - 6 R. Takasawa, S. Takahashi, K. Saeki, S. Sunaga, A. Yoshimoria and S. Tanumaa, *Bioorg. Med. Chem. Lett.*, 2008, **16**, 3969–3975.
 - 7 M. Kawatani, H. Okumura, K. Honda, N. Kanoh, M. Muroi,
 N. Dohmae, M. Takami, M. Kitagawa, Y. Futamura,
 M. Imoto and H. Osada, *Proc. Natl. Acad. Sci. U. S. A.*, 2008,
 105, 11691–11696.
- 8 R. Takasawa, K. Saeki, A. Tao, A. Yoshimori, H. Uchiro, M. Fujiwara and S. Tanuma, *Bioorg. Med. Chem. Lett.*, 2010, 18, 7029–7033.
 - 9 M. Liu, M. Yuan, M. Luo, X. Bu, H. Luo and X. Hu, *Biophys. Chem.*, 2010, 147, 28–34.
- 55 10 J. Zhai, H. Zhang, L. Zhang, Y. Zhao, S. Chen, Y. Chen, X. Peng, Q. Li, M. Yuan and X. Hu, *ChemMedChem*, 2013, 8, 1462–1464.
 - 11 R. Nelson and A. Rosowsky, Antimicrob. Agents Chemother., 2001, 45, 3293–3303.

- 12 A. Zhao, X. Gao, Y. Wang, J. Ai, Y. Wang, Y. Chen, M. Geng and A. Zhang, *Bioorg. Med. Chem.*, 2011, **19**, 3906–3918.
- M. Argiriadi, A. Ericsson, C. Harris, D. Banach, D. Borhani,
 D. Calderwood, M. Demers, J. Dimauro, R. Dixon,
 J. Hardman, S. Kwak, B. Li, J. Mankovich, D. Marcotte,
 K. Mullen, B. Ni, M. Pietras, R. Sadhukhan, S. Sousa,
 M. Tomlinson, L. Wang, T. Xiang and R. Talanian, *Bioorg. Med. Chem. Lett.*, 2010, 20, 330–333.
- 14 C. Zificsak, J. Theroff, L. Aimone, T. Angeles, M. Albom, M. Cheng, E. Mesaros, G. Ott, M. Quail, T. Underiner, 10
 W. Wan and B. Dorsey, *Bioorg. Med. Chem. Lett.*, 2011, 21, 3877–3880.
- 15 N. Powell, J. Hoffman, F. Ciske, M. Kaufman, J. Kohrt, J. Quin, D. Sheehan, A. Delaney, S. Baxi, C. Catana, P. McConnell, J. Ohren, L. Perrin and J. Edmunds, *Bioorg.* 15 *Med. Chem. Lett.*, 2013, 23, 1046–1050.
- 16 N. Ramsden, J. Perrin, Z. Ren, B. Lee, N. Zinn, V. Dawson, D. Tam, M. Bova, M. Lang, G. Drewes, M. Bantscheff, F. Bard, T. Dawson and C. Hopf, ACS Chem. Biol., 2011, 6, 1021–1028.
- 17 H. Chen, B. Chan, J. Drummond, A. Estrada, J. Gunzner-Toste, X. Liu, Y. Liu, J. Moffat, D. Shore, Z. Sweeney, T. Tran, S. Wang, G. Zhao, H. Zhu and D. Burdick, *J. Med. Chem.*, 2012, **14**, 5536–5545.
- 18 H. Choi, J. Zhang, X. Deng, J. Hatcher, M. Patricelli, Z. Zhao, D. Alessi and N. Gray, *ACS Med. Chem. Lett.*, 2012, **3**, 658–662.
- 19 J. Bain, L. Plater, M. Elliott, N. Shpiro, C. J. Hastie, H. McLauchlan, I. Klevernic, J. S. Arthur, D. R. Alessi and P. Cohen, *Biochem. J.*, 2007, 408, 297–315.
- 20 M. W. Karaman, S. Herrgard, D. K. Treiber, P. Gallant, C. E. Atteridge, B. T. Campbell, K. W. Chan, P. Ciceri, M. I. Davis, P. T. Edeen, R. Faraoni, M. Floyd, J. P. Hunt, D. J. Lockhart, Z. V. Milanov, M. J. Morrison, G. Pallares, H. K. Patel, S. Pritchard, L. M. Wodicka and 35 P. P. Zarrinkar, *Nat. Biotechnol.*, 2008, 26, 127–132.
- 21 M. P. Patricelli, T. K. Nomanbhoy, J. Wu, H. Brown, D. Zhou,
 J. Zhang, S. Jagannathan, A. Aban, E. Okerberg, C. Herring,
 B. Nordin, H. Weissig, Q. Yang, J. D. Lee, N. S. Gray and
 J. W. Kozarich, *Chem. Biol.*, 2011, 18, 699–710.
- 22 M. Schenone, V. Dancik, B. K. Wagner and P. A. Clemons, *Nat. Chem. Biol.*, 2013, **9**, 232–240.
- 23 A. Cameron, B. Olin, M. Ridderström, B. Mannervik and T. Jones, *EMBO J.*, 1997, **16**, 3386–3395.
- 45
 24 A. Speers and B. Cravatt, Profiling enzyme activities *in vivo* using click chemistry methods, *Chem. Biol.*, 2004, **11**, 535–546.
- 25 P. Thornalley, Biochem. J., 1988, 254, 751-755.
- 26 M. Shinohara, P. Thornalley, I. Giardino, P. Beisswenger, 50
 S. Thorpe, J. Onorato and M. Brownlee, *J. Clin. Invest.*, 1998, 101, 1142–1147.
- 27 H. Sakamoto, T. Mashima, S. Sato, Y. Hashimoto, T. Yamori and T. Tsuruo, *Clin. Cancer Res.*, 2001, 7, 2513–2518.
- 28 N. Shangari, W. Bruce, R. Poon and P. O'Brien, *Biochem. Soc.* 55 *Trans.*, 2003, 31, 1390–1393.
- 29 O. Warburg, Science, 1956, 123, 309-314.
- 30 R. Gatenby and R. Gillies, Nat. Rev. Cancer, 2004, 4, 891–899.

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	MedChemComm	Concise Article	
1	31 A. Rulli, L. Carli, R. Romani, T. Baroni, E. Giovannini, G. Rosi and V. Talesa, <i>Breast Cancer Res. Treat.</i> , 2001, 66 , 67–72.	32 C. Antognelli, F. Baldracchini, V. Talesa, E. Costantini, A. Zucchi and E. Mearini, <i>Cancer J.</i> , 2006, 12 , 222–228.	1
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